

SPECIFICATION AMENDMENTS

(Insertions indicated by underline; deletions indicated by strikethrough)

Please replace the paragraph beginning at page 43, line 25, with:

A
In order to identify novel Ras-family gene products which have tumor suppressor activity, an electronic screen was undertaken to identify proteins showing homology to the Noey2 (also called Ahril) gene product (Yu *et al.*, *Proc. Natl. Acad. Sci. USA* 96:214-219 [1999]; GenBank Accession No. NP_004666; and SEQ ID NO: 9). The full length Noey2 amino acid sequence (SEQ ID NO: 9) was used as the query sequence to search the databases of the National Center for Biotechnology Information (NCBI), using the search program "Advanced tBLASTn" (><http://www.ncbi.nlm.nih.gov/BLAST/><). This electronic screen identified an open reading frame (ORF) on bacterial artificial chromosome (BAG) 41195 (GenBank Accession No. AC006538), corresponding to a region on human chromosome 19 at pi3.3. This ORF was then used to search the NCBI expressed sequence tag (EST) databases. That search identified an IMAGE Consortium EST (IMAGE Consortium Clone ID No. 2165313; GenBank Accession No. AI497811) containing a human ORF corresponding to the region of interest on the BAG 41195 clone. This IMAGE Consortium Clone was obtained from ATCC (Catalog No. 3363561) and used as template material in a PCR strategy to isolate the full length ORF. These primers contained terminal *Bam*HI and *Eco*RI restriction sites, respectively, to facilitate subcloning. The resulting PCR product was subcloned into a variety of vectors for subsequent study.